



SEQUENCE LISTING

<110> NASU, Hisanori
ONO, Hiroaki
MUGITA, Akito
NAKAYAMA, Ichiro
KOBAYASHI, Takanori
MASAOKA, Tetsuji
KUHARA, Satoru
ISHINO, Yoshizumi

<120> METHOD FOR DETECTION OF BASE SEQUENCE OF INTEREST

<130> Q-80511

<140> US 10/802,055

<141> 2004-03-17

<160> 6

<170> PatentIn version 3.3

<210> 1

<211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Target Sequence

<400> 1

accagaaaaa atttatTTTT tctggtgttt gccgcccggc tccagtggaa atgaaactct 60

ggg 63

<210> 2

<211> 16

<212> DNA

<213> Artificial Sequence

<220>

<223> Probe

<400> 2

gcggcaaaca ccagaa 16

<210> 3

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> Probe

<400> 3
ttcatttcca ctgga 15

<210> 4
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Probe

<400> 4
gccaacatat gtcctctgac gc 22

<210> 5
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Probe

<400> 5
cgcttaaggt tactaacctg tcgagg 26

<210> 6
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Probe

<400> 6
gcgaacatat ggtcgcctct atcga 25